

Figure 1.

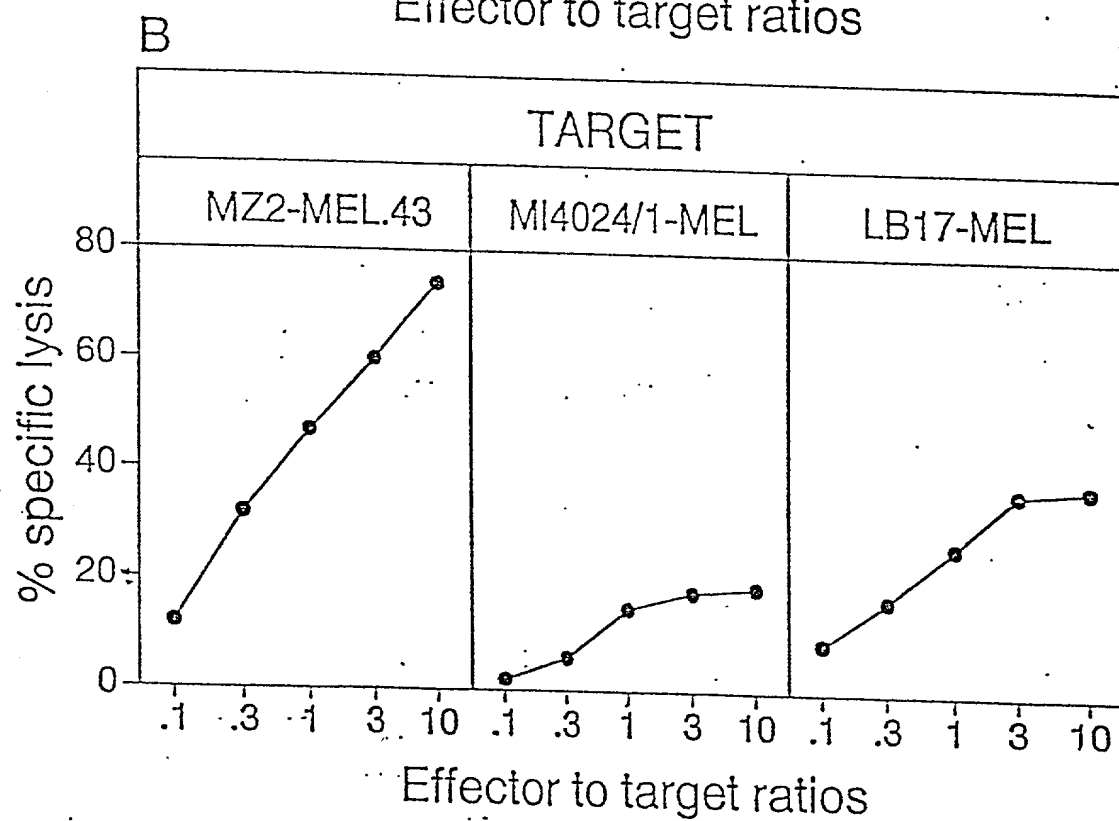
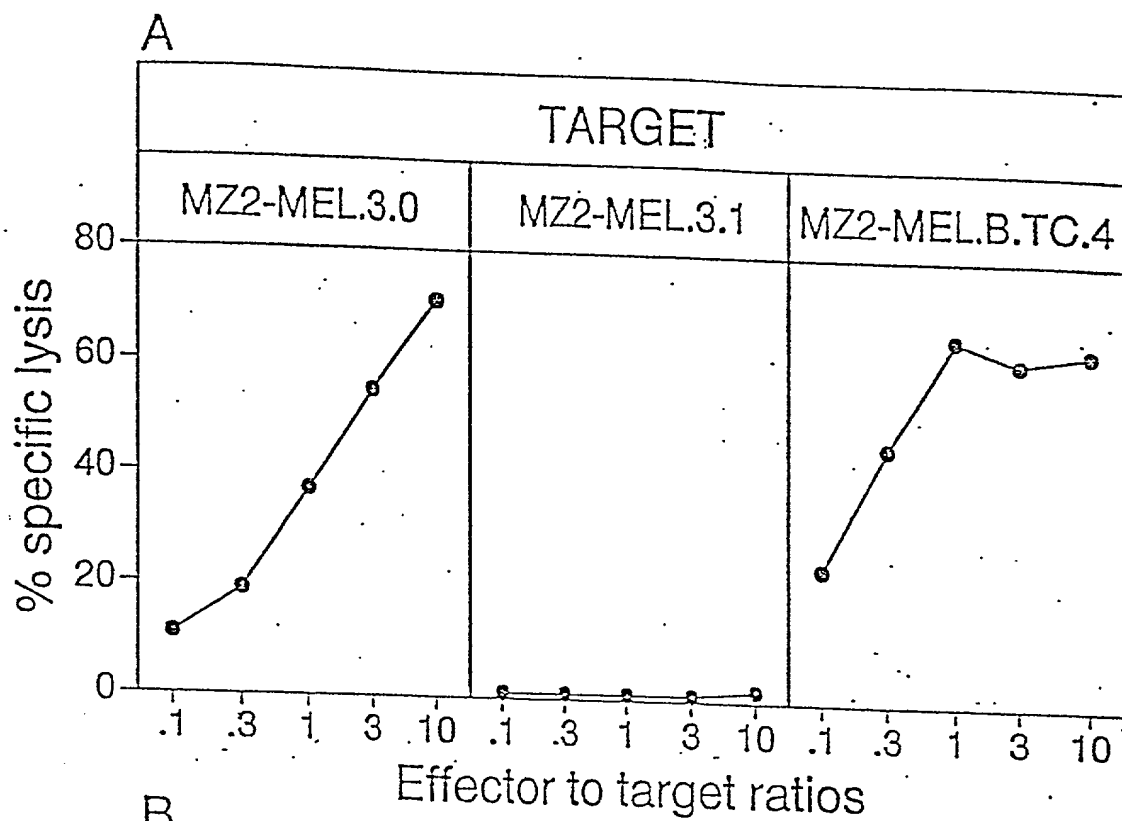


Figure 3

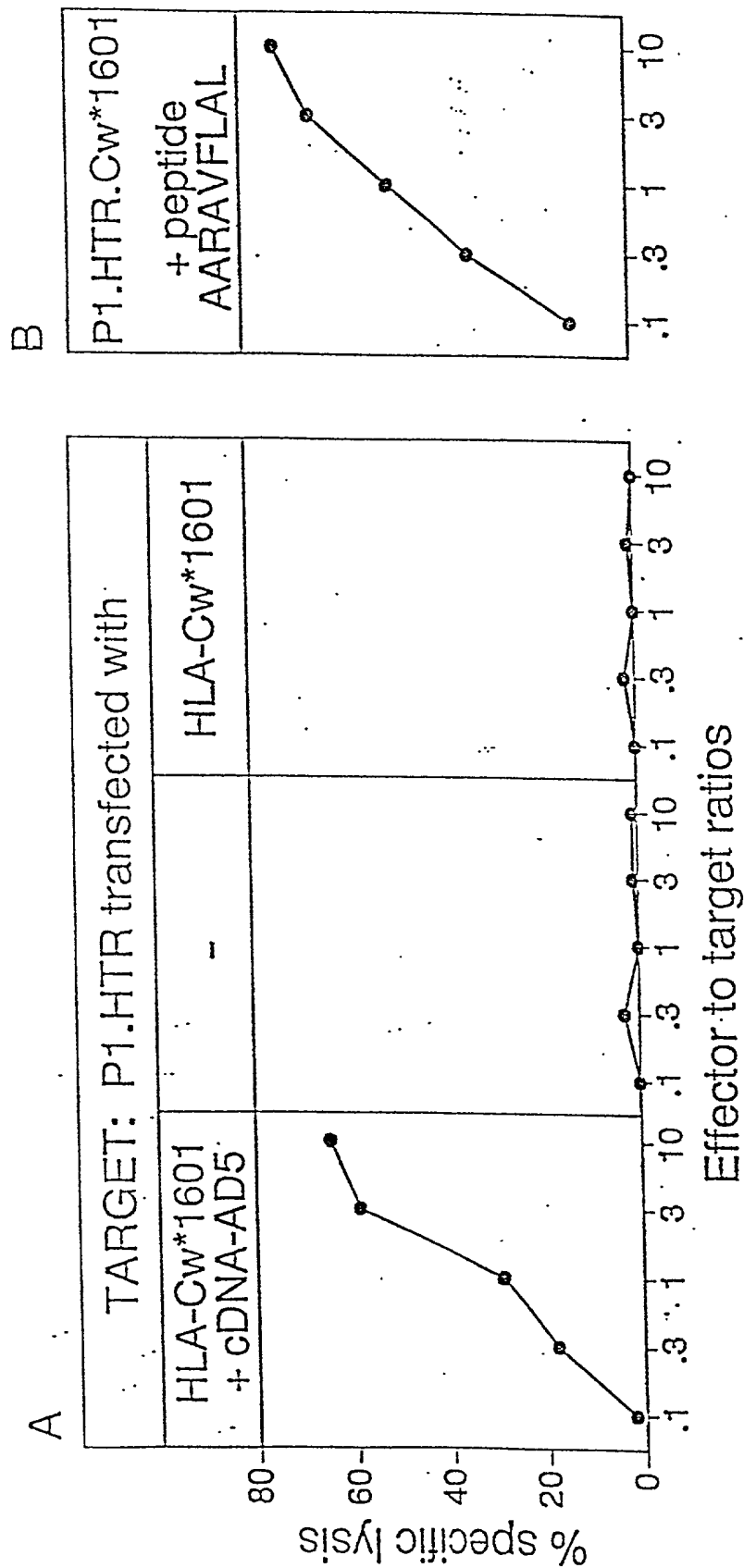


Figure 4

CGCCAATTTA	GGGTCTCCGG	TATCTCCCGC	TGAGCTGCTC	TGTTCCCGGC	TTAGAGGACC	60									
AGGAGAAGGG	GGAGCTGGAG	GCTGGAGCCT	GTAACACCGT	GGCTCGTCTC	ACTCTGGATG	120									
GTGGTGGCAA	CAGAGATGGC	AGCGCAGCTG	GAGTGTTAGG	AGGGCGGCCT	GAGCGGTAGG	180									
		M	A	A	R	A	V	F	L	A	L	S	A	Q	13
AGTGGGGCTG	GAGCAGTAAG	ATGGCGGCCA	GAGCGGTTTT	TCTGGCATTG	TCTGCCCAGC	240									
L L Q A	R L M	K E E	S P V V	S W R	L E P	33									
TGCTCCAAGC	CAGGCTGATG	AAGGAGGAGT	CCCCTGTGGT	GAGCTGGAGG	TTGGAGCCTG	300									
E D G T	A L C	F I F				43									
AAGACGGCAC	AGCTCTGTGC	TTCATCTTCT	GAGGTTGTGG	CAGCCACGGT	GATGGAGACG	360									
GCAGCTCAAC	AGGAGCAATA	GGAGGAGATG	GAGTTTCACT	GTGTCAGCCA	GGATGGTCTC	420									
GATCTCCTGA	CCTCGTGATC	CGCCCGCCTT	GGCCTTCCAA	AGTGCCGAGA	TTACAGCGAT	480									
GTGCATTTTG	TAAGCACTTT	GGAGCCACTA	TCAAATGCTG	TGAAGAGAAA	TGTACCCAGA	540									
TGTATCATT	TCCTTGTGCT	GCAGGAGCCG	GCTCCTTTCA	GGATTTCACT	CACATCTTCC	600									
TGCTTTGTCC	AGAACACATT	GACCAAGCTC	CTGAAAGATG	TAAGTTTACT	ACGCATAGAC	660									
TTTATAACTT	CAACCAATGT	ATTTACTGAA	AATAACAAAT	GTTGTAAATT	CCCTGAGTGT	730									
TATCTTACTT	GTATTAAAAG	GTAATAATAC	ATAATCATT	AAATCTGAGG	GATCATTGCC	780									
AGAGATTGTT	GGGGAGGGAA	ATGTTATCAA	CGGTTTCATT	GAAATTAAAT	CCAAAAAGTT	840									
ATTCCTCAG	AAAAATCAAA	TAAAGTTTGC	ATGTTTTTTA	TTCTTAAAAC	ATTTTAAAAA	900									
CCACTGTAGA	ATGATGTAAA	TAGGGACTGT	GCAGTATTTT	TGACATATAC	TATAAAATTA	960									
TTAAAAAGTC	AATCAGTATT	CAACATCTTT	TACACTAAAA	AGCC		1004									

Figure 5

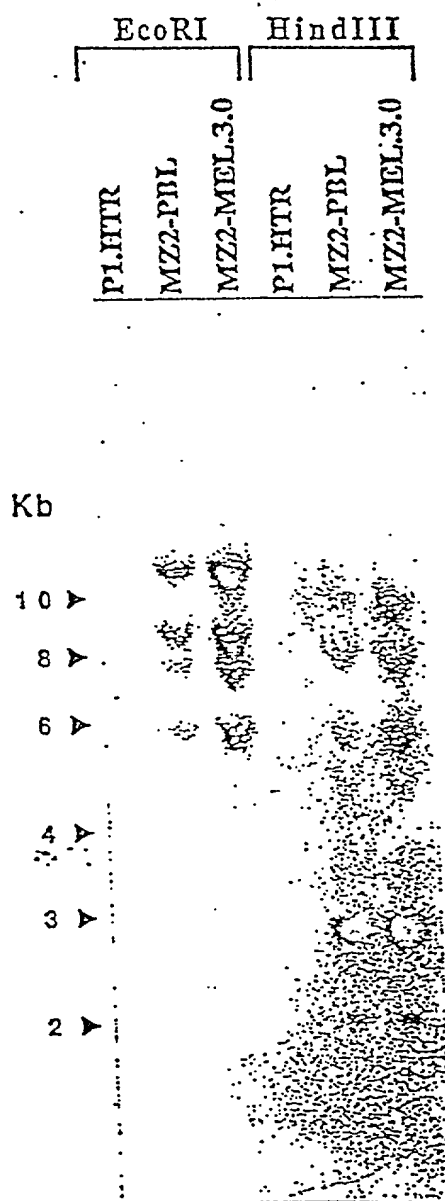


Figure 6

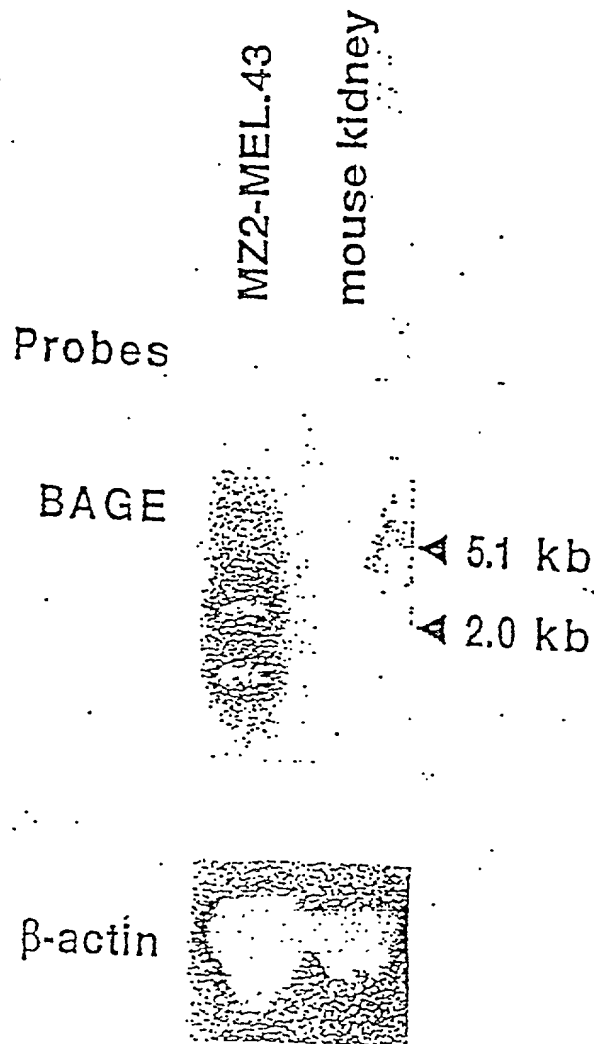


Figure 7

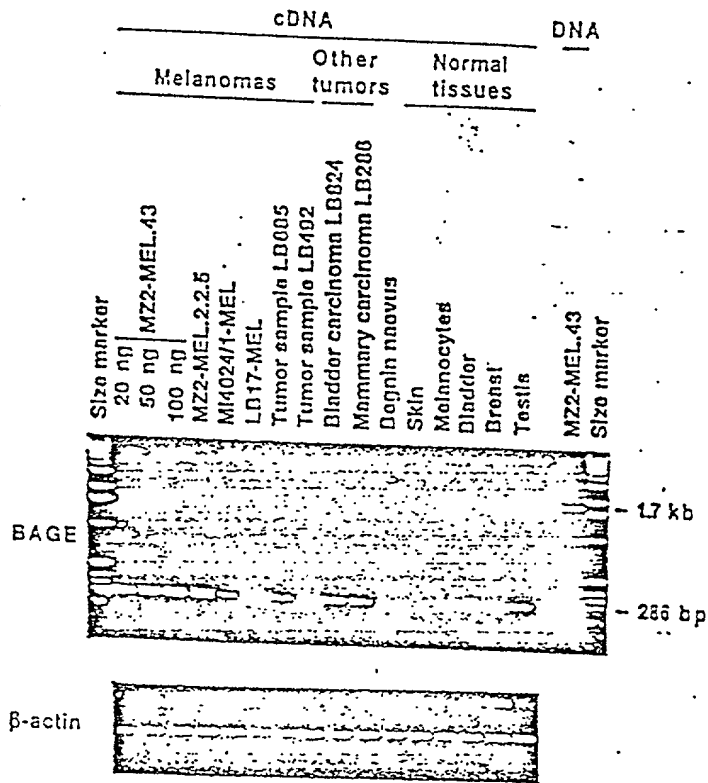


Figure 8

